

# SEQUENCE LISTING

<110> Yale University

<120> TREATMENT AND PREVENTION OF CANCER AND PITUITARY  
DISORDERS WITH LATS PROTEINS, DERIVATIVES AND  
FRAGMENTS, AND LATS KNOCK-OUT ANIMAL MODELS

<130> 6523-020-999

<140> 09/763,334

<141> 2001-02-16

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<170> PatentIn Ver. 2.0

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Ser Leu Val Pro Gln Arg His Gly Pro Ser Leu Gly Glu Asn Val Val	
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Tyr Arg Ser Glu Ser Pro Asn Ser Gln Ala Asp Val Gly Arg Pro Leu	
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Gln Arg Val Asn Pro Pro Pro Pro Gln Val Arg Ser Val Thr Pro	
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Asn Val Pro Gly Leu Gln Thr Ala Trp Pro Gln Ser Ser Ser Ala Pro	
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Ser Asp Leu Arg Gln Gln Ser Ala Ser Tyr Ile Pro Lys Ile Thr His	
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Pro Thr Asp Thr Ser Asn Phe Asp Pro Val Asp Pro Asp Lys Leu Trp	
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Tyr Lys Asn Gly Lys His Pro Glu His Ala Phe Tyr Glu Phe Thr Phe	
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 Ala Ser Ser Val Pro Val Ile Pro Pro Val Ala Glu Ala Pro Ser Tyr  
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 Pro Ser Leu Pro Lys Glu Asp Asp Ser Glu Lys Ser Ala Asp Ser Gly  
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Cys	Leu	Ala	His	Ser	Leu	Val	Gly	Thr	Pro	Asn	Tyr	Ile	Ala	Pro	Glu	
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Thr Gly Glu Ala Leu Pro Ser Tyr His Gln Leu Gly Gly Ala Asn Tyr	
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Lys Pro Pro Ser Cys Asn Asn Asn Asn Ile Gln Ile Ser Asn Ser Asn	
555 560 565 570	
ctg gcg acg aca cca ccc att ccg cct gcc aaa tac aat aac aac tcc	2860
Leu Ala Thr Thr Pro Pro Ile Pro Pro Ala Lys Tyr Asn Asn Asn Ser	
575 580 585	
tcc aac acg ggc gcg aat agc tcg ggc ggc agc aac gga tcc acc ggc	2908
Ser Asn Thr Gly Ala Asn Ser Ser Gly Gly Ser Asn Gly Ser Thr Gly	
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acc acc gcc tcc tcg tcg acc agc tgc aag aag atc aag cac gcc tcg	2956
Thr Thr Ala Ser Ser Ser Thr Ser Cys Lys Lys Ile Lys His Ala Ser	
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ccc atc ccg gag cgc aag aag atc tcc aag gag aag gag gag gag cgc	3004
Pro Ile Pro Glu Arg Lys Lys Ile Ser Lys Glu Lys Glu Glu Glu Arg	
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aag gag ttc cgc atc agg cag tac tcg ccg caa gcc ttc aag ttc ttc	3052
Lys Glu Phe Arg Ile Arg Gln Tyr Ser Pro Gln Ala Phe Lys Phe Phe	
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atg gag cag cac ata gag aac gtg atc aag tcg tat cgc cag cgc acg	3100
Met Glu Gln His Ile Glu Asn Val Ile Lys Ser Tyr Arg Gln Arg Thr	
655 660 665	
tat cgc aag aat cag ctg gag aag gag atg cac aaa gtg gga ctg ccc	3148
Tyr Arg Lys Asn Gln Leu Glu Lys Glu Met His Lys Val Gly Leu Pro	
670 675 680	
gat cag acc caa atc gag atg agg aaa atg ctg aac caa aag gag agc	3196
Asp Gln Thr Gln Ile Glu Met Arg Lys Met Leu Asn Gln Lys Glu Ser	
685 690 695	

aac tac att cga ttg aag cgc gcc aag atg gac aag agc atg ttc gtc	3244
Asn Tyr Ile Arg Leu Lys Arg Ala Lys Met Asp Lys Ser Met Phe Val	
700 705 710	
aaa ctg aag ccc att gga gtg ggt gca ttt ggc gag gta acg ctg gtg	3292
Lys Leu Lys Pro Ile Gly Val Gly Ala Phe Gly Glu Val Thr Leu Val	
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agc aaa atc gat acc tcg aac cat ttg tat gcg atg aaa acc ctg cgg	3340
Ser Lys Ile Asp Thr Ser Asn His Leu Tyr Ala Met Lys Thr Leu Arg	
735 740 745	
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Lys Ala Asp Val Leu Lys Arg Asn Gln Val Ala His Val Lys Ala Glu	
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agg gat atc ctc gcg gaa gcc gac aat aac tgg gtg gtg aag ttg tac	3436
Arg Asp Ile Leu Ala Glu Ala Asp Asn Asn Trp Val Val Lys Leu Tyr	
765 770 775	
tac agc ttc cag gac aag gat aat ctg tac ttt gtg atg gac tac ata	3484
Tyr Ser Phe Gln Asp Lys Asp Asn Leu Tyr Phe Val Met Asp Tyr Ile	
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cca ggt ggt gat ctg atg tcg ctg ctc atc aaa ctg ggc att ttc gag	3532
Pro Gly Gly Asp Leu Met Ser Leu Leu Ile Lys Leu Gly Ile Phe Glu	
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gag gaa ctg gcc aga ttc tac atc gcc gag gtc acc tgc gcc gtg gac	3580
Glu Glu Leu Ala Arg Phe Tyr Ile Ala Glu Val Thr Cys Ala Val Asp	
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agc gtt cac aaa atg ggc ttc att cac aga gac atc aag cct gac aac	3628
Ser Val His Lys Met Gly Phe Ile His Arg Asp Ile Lys Pro Asp Asn	
830 835 840	
ata ctc atc gat agg gac gga cac ata aag ctc acc gac ttt ggc ctg	3676
Ile Leu Ile Asp Arg Asp Gly His Ile Lys Leu Thr Asp Phe Gly Leu	
845 850 855	
tgc acg gga ttc cga tgg acg cac aac tcg aag tac tac cag gag aac	3724
Cys Thr Gly Phe Arg Trp Thr His Asn Ser Lys Tyr Tyr Gln Glu Asn	
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ggc aat cac tcg cgc cag gac tcg atg gag ccc tgg gag gaa tac tcc	3772
Gly Asn His Ser Arg Gln Asp Ser Met Glu Pro Trp Glu Glu Tyr Ser	
875 880 885 890	
gag aac gga ccg aag ccc acc gtg ctg gag agg cga cgg atg cgc gat	3820
Glu Asn Gly Pro Lys Pro Thr Val Leu Glu Arg Arg Arg Met Arg Asp	
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His Gln Arg Val Leu Ala His Ser Leu Val Gly Thr Pro Asn Tyr Ile	
910 915 920	
gct ccc gag gtg ctg gag agr agt ggg tac acg cag ctg tgc gac tac	3916
Ala Pro Glu Val Leu Glu Arg Ser Gly Tyr Thr Gln Leu Cys Asp Tyr	
925 930 935	
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Trp Ser Val Gly Val Ile Leu Tyr Glu Met Leu Val Gly Gln Pro Pro	
940 945 950	
ttt ctg gcc aac agt ccg ctg gaa acg caa caa aag gtc atc aac tgg	4012
Phe Leu Ala Asn Ser Pro Leu Glu Thr Gln Gln Lys Val Ile Asn Trp	
955 960 965 970	
gag aaa acs ctg cat att ccg ccg cag gcc gag tta tcc cgc gag gct	4060
Glu Lys Thr Leu His Ile Pro Pro Gln Ala Glu Leu Ser Arg Glu Ala	
975 980 985	
acg gac ttg ata agg agg ctc tgt gcg tcg gct gac aag cgg ctg ggc	4108
Thr Asp Leu Ile Arg Arg Leu Cys Ala Ser Ala Asp Lys Arg Leu Gly	
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Lys Ser Val Asp Glu Val Lys Ser His Asp Phe Phe Lys Gly Ile	
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Asp Phe Ala Asp Met Arg Lys Gln Lys Ala Pro Tyr Ile Pro Glu	
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Ile Lys His Pro Thr Asp Thr Ser Asn Phe Asp Pro Val Asp Pro	
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Glu Lys Leu Arg Ser Asn Asp Ser Thr Met Ser Ser Gly Asp Asp	
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Val Asp Gln Asn Asp Arg Thr Phe His Gly Phe Phe Glu Phe Thr	
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Phe Arg Arg Phe Phe Asp Asp Lys Gln Pro Pro Asp Met Thr Asp	
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Asp Gln Ala Pro Val Tyr Val	
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 <213> *Drosophila melanogaster*

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 <223> Xaa = Lys, Ser

<220>  
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Val	Gln	Asn	Asn	His	Arg	Asn	Asn	Gln	Asn	Tyr	Thr	Pro	Leu	Arg	Tyr
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Val	Val	Ile	Pro	Pro	Pro	Pro	Ala	Ile	Val	Gly	Gln	Pro	Gly	Ala	Gly
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Ser	Ile	Ser	Val	Ser	Gly	Val	Gly	Val	Gly	Val	Val	Gly	Val	Ala	Asn
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Gly	Arg	Val	Pro	Lys	Met	Met	Thr	Ala	Leu	Met	Pro	Asn	Lys	Leu	Ile
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Arg	Lys	Pro	Ser	Ile	Glu	Arg	Asp	Thr	Ala	Ser	Ser	His	Tyr	Leu	Arg
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Cys	Ser	Pro	Ala	Leu	Asp	Ser	Gly	Ala	Gly	Ser	Ser	Arg	Ser	Asp	Ser
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Pro	Gly	Gly	Asn	Gly	Gly	Phe	Ser	Pro	Ser	Pro	Ser	Gly	Phe	Ser	Glu
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Val	Ala	Pro	Pro	Ala	Pro	Pro	Pro	Arg	Asn	Pro	Thr	Ala	Xaa	Ser	Ala
		195					200					205			
Ala	Thr	Pro	Pro	Pro	Pro	Val	Pro	Pro	Thr	Ser	Gln	Ala	Tyr	Val	Lys
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Arg	Arg	Ser	Pro	Ala	Leu	Asn	Asn	Arg	Pro	Pro	Ala	Ile	Ala	Pro	Pro
225					230					235					240
Thr	Gln	Arg	Gly	Asn	Ser	Pro	Val	Ile	Thr	Gln	Asn	Gly	Leu	Lys	Asn
				245					250					255	
Pro	Gln	Gln	Gln	Leu	Thr	Gln	Gln	Leu	Lys	Ser	Leu	Asn	Leu	Tyr	Pro
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Gly	Gly	Gly	Ser	Gly	Ala	Val	Val	Glu	Pro	Pro	Pro	Pro	Tyr	Leu	Ile
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Gln	Gly	Gly	Ala	Gly	Gly	Ala	Ala	Pro	Pro	Pro	Pro	Pro	Pro	Ser	Tyr
	290					295						300			

Thr Ala Ser Met Gln Ser Arg Gln Ser Pro Thr Gln Ser Gln Gln Ser  
 305 310 315 320  
 Asp Tyr Arg Lys Ser Pro Ser Ser Gly Ile Tyr Ser Ala Thr Ser Ala  
 325 330 335  
 Gly Ser Pro Ser Pro Ile Thr Val Ser Leu Pro Pro Ala Pro Leu Ala  
 340 345 350  
 Lys Pro Gln Pro Arg Val Tyr Gln Ala Arg Ser Gln Gln Pro Ile Ile  
 355 360 365  
 Met Gln Ser Val Lys Ser Thr Gln Val Gln Lys Pro Val Leu Gln Thr  
 370 375 380  
 Ala Val Ala Arg Gln Ser Pro Ser Ser Ala Ser Ala Ser Asn Ser Pro  
 385 390 395 400  
 Val His Val Leu Ala Ala Pro Pro Ser Tyr Pro Gln Lys Ser Ala Ala  
 405 410 415  
 Val Val Gln Gln Gln Gln Gln Ala Ala Ala Ala His Gln Gln Gln  
 420 425 430  
 His Gln His Gln Gln Ser Lys Pro Pro Thr Pro Thr Thr Pro Pro Leu  
 435 440 445  
 Val Gly Leu Asn Ser Lys Pro Asn Cys Leu Glu Pro Pro Ser Tyr Ala  
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 Lys Ser Met Gln Ala Lys Ala Ala Thr Val Val Xaa Gln Gln Gln Gln  
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 Gln Gln Gln Gln Gln Gln Leu Gln Ala Leu Arg Val Leu Gln Ala Gln  
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 Ala Gln Arg Glu Arg Asp Gln Arg Glu Arg Glu Arg Asp Gln Gln Lys  
 515 520 525  
 Leu Ala Asn Gly Asn Pro Gly Arg Gln Met Leu Pro Pro Pro Pro Tyr  
 530 535 540  
 Gln Ser Asn Asn Asn Asn Asn Ser Glu Ile Lys Pro Pro Ser Cys Asn  
 545 550 555 560  
 Asn Asn Asn Ile Gln Ile Ser Asn Ser Asn Leu Ala Thr Thr Pro Pro  
 565 570 575  
 Ile Pro Pro Ala Lys Tyr Asn Asn Asn Ser Ser Asn Thr Gly Ala Asn  
 580 585 590  
 Ser Ser Gly Gly Ser Asn Gly Ser Thr Gly Thr Thr Ala Ser Ser Ser  
 595 600 605

Thr Ser Cys Lys Lys Ile Lys His Ala Ser Pro Ile Pro Glu Arg Lys  
610 615 620

Lys Ile Ser Lys Glu Lys Glu Glu Glu Arg Lys Glu Phe Arg Ile Arg  
625 630 635 640

Gln Tyr Ser Pro Gln Ala Phe Lys Phe Phe Met Glu Gln His Ile Glu  
645 650 655

Asn Val Ile Lys Ser Tyr Arg Gln Arg Thr Tyr Arg Lys Asn Gln Leu  
660 665 670

Glu Lys Glu Met His Lys Val Gly Leu Pro Asp Gln Thr Gln Ile Glu  
675 680 685

Met Arg Lys Met Leu Asn Gln Lys Glu Ser Asn Tyr Ile Arg Leu Lys  
690 695 700

Arg Ala Lys Met Asp Lys Ser Met Phe Val Lys Leu Lys Pro Ile Gly  
705 710 715 720

Val Gly Ala Phe Gly Glu Val Thr Leu Val Ser Lys Ile Asp Thr Ser  
725 730 735

Asn His Leu Tyr Ala Met Lys Thr Leu Arg Lys Ala Asp Val Leu Lys  
740 745 750

Arg Asn Gln Val Ala His Val Lys Ala Glu Arg Asp Ile Leu Ala Glu  
755 760 765

Ala Asp Asn Asn Trp Val Val Lys Leu Tyr Tyr Ser Phe Gln Asp Lys  
770 775 780

Asp Asn Leu Tyr Phe Val Met Asp Tyr Ile Pro Gly Gly Asp Leu Met  
785 790 795 800

Ser Leu Leu Ile Lys Leu Gly Ile Phe Glu Glu Glu Leu Ala Arg Phe  
805 810 815

Tyr Ile Ala Glu Val Thr Cys Ala Val Asp Ser Val His Lys Met Gly  
820 825 830

Phe Ile His Arg Asp Ile Lys Pro Asp Asn Ile Leu Ile Asp Arg Asp  
835 840 845

Gly His Ile Lys Leu Thr Asp Phe Gly Leu Cys Thr Gly Phe Arg Trp  
850 855 860

Thr His Asn Ser Lys Tyr Tyr Gln Glu Asn Gly Asn His Ser Arg Gln  
865 870 875 880

Asp Ser Met Glu Pro Trp Glu Glu Tyr Ser Glu Asn Gly Pro Lys Pro  
885 890 895

Thr Val Leu Glu Arg Arg Arg Met Arg Asp His Gln Arg Val Leu Ala  
900 905 910

His Ser Leu Val Gly Thr Pro Asn Tyr Ile Ala Pro Glu Val Leu Glu  
 915 920 925  
 Arg Ser Gly Tyr Thr Gln Leu Cys Asp Tyr Trp Ser Val Gly Val Ile  
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 Xaa Tyr Glu Met Leu Val Gly Gln Pro Pro Phe Leu Ala Asn Ser Pro  
 945 950 955 960  
 Leu Glu Thr Gln Gln Lys Val Ile Asn Trp Glu Lys Thr Leu His Ile  
 965 970 975  
 Pro Pro Gln Ala Glu Leu Ser Arg Glu Ala Thr Asp Leu Ile Arg Arg  
 980 985 990  
 Leu Cys Ala Ser Ala Asp Lys Arg Leu Gly Lys Ser Val Asp Glu Val  
 995 1000 1005  
 Lys Ser His Asp Phe Phe Lys Gly Ile Asp Phe Ala Asp Met Arg  
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 Lys Gln Lys Ala Pro Tyr Ile Pro Glu Ile Lys His Pro Thr Asp  
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 Thr Ser Asn Phe Asp Pro Val Asp Pro Glu Lys Leu Arg Ser Asn  
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 Asp Ser Thr Met Ser Ser Gly Asp Asp Val Asp Gln Asn Asp Arg  
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 Thr Phe His Gly Phe Phe Glu Phe Thr Phe Arg Arg Phe Phe Asp  
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 Asp Lys Gln Pro Pro Asp Met Thr Asp Asp Gln Ala Pro Val Tyr  
 1085 1090 1095

Val

<210> 9  
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 <212> PRT  
 <213> Mus musculus

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 Val Tyr Arg Ser Glu Ser Pro Asn Ser Gln Ala Asp Val Gly Arg Pro  
 35 40 45  
 Leu Ser Gly Ser Gly Ile Ala Ala Phe Ala Gln Ala His Pro Ser Asn  
 50 55 60  
 Gly Gln Arg Val Asn Pro Pro Pro Pro Gln Val Arg Ser Val Thr  
 65 70 75 80  
 Pro Pro Pro Pro Pro Arg Gly Gln Thr Pro Pro Pro Arg Gly Thr Thr  
 85 90 95

Pro	Pro	Pro	Pro	Ser	Trp	Glu	Pro	Ser	Ser	Gln	Thr	Lys	Arg	Tyr	Ser
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Gly	Asn	Met	Glu	Tyr	Val	Ile	Ser	Arg	Ile	Ser	Pro	Val	Pro	Pro	Gly
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Ala	Trp	Gln	Glu	Gly	Tyr	Pro	Pro	Pro	Pro	Leu	Thr	Thr	Ser	Pro	Met
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Asn	Pro	Pro	Ser	Gln	Ala	Gln	Arg	Ala	Ile	Ser	Ser	Val	Pro	Val	Gly
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Arg	Gln	Pro	Ile	Ile	Met	Gln	Ser	Thr	Ser	Lys	Phe	Asn	Phe	Thr	Pro
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Gly	Arg	Pro	Gly	Val	Gln	Asn	Gly	Gly	Gly	Gln	Ser	Asp	Phe	Ile	Val
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His	Gln	Asn	Val	Pro	Thr	Gly	Ser	Val	Thr	Arg	Gln	Pro	Pro	Pro	Pro
		195					200					205			
Tyr	Pro	Leu	Thr	Pro	Ala	Asn	Gly	Gln	Ser	Pro	Ser	Ala	Leu	Gln	Thr
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Gly	Ala	Ser	Ala	Ala	Pro	Pro	Ser	Phe	Ala	Asn	Gly	Asn	Val	Pro	Gln
225					230					235					240
Ser	Met	Met	Val	Pro	Asn	Arg	Asn	Ser	His	Asn	Met	Glu	Leu	Tyr	Asn
			245						250					255	
Ile	Asn	Val	Pro	Gly	Leu	Gln	Thr	Ala	Trp	Pro	Gln	Ser	Ser	Ser	Ala
		260						265					270		
Pro	Ala	Gln	Ser	Ser	Pro	Ser	Gly	Gly	His	Glu	Ile	Pro	Thr	Trp	Gln
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Arg	Ala	Ser	His	Ser	Ala	Asn	Ser	Gln	Pro	Ser	Ala	Thr	Thr	Val	Thr
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Ala	Ile	Thr	Pro	Ala	Pro	Ile	Gln	Gln	Pro	Val	Lys	Ser	Met	Arg	Val
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Leu	Lys	Pro	Glu	Leu	Gln	Thr	Ala	Leu	Ala	Pro	Thr	His	Pro	Ser	Trp
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Met	Pro	Gln	Pro	Val	Gln	Thr	Val	Gln	Pro	Thr	Pro	Phe	Ser	Glu	Gly
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Thr	Ala	Ser	Ser	Val	Pro	Val	Ile	Pro	Pro	Val	Ala	Glu	Ala	Pro	Ser
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Tyr	Gln	Gly	Pro	Pro	Pro	Pro	Tyr	Pro	Lys	His	Leu	Leu	His	Gln	Asn
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Pro	Ser	Val	Pro	Pro	Tyr	Glu	Ser	Val	Ser	Lys	Pro	Cys	Lys	Asp	Glu
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Gln	Pro	Ser	Leu	Pro	Lys	Glu	Asp	Asp	Ser	Glu	Lys	Ser	Ala	Asp	Ser
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Gly	Asp	Ser	Gly	Asp	Lys	Glu	Lys	Lys	Gln	Ile	Thr	Thr	Ser	Pro	Ile
	435						440					445			
Thr	Val	Arg	Lys	Asn	Lys	Lys	Asp	Glu	Glu	Arg	Arg	Glu	Ser	Arg	Ile
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Gln	Ser	Tyr	Ser	Pro	Gln	Ala	Phe	Lys	Phe	Phe	Met	Glu	Gln	His	Val
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Glu	Asn	Val	Leu	Lys	Ser	His	Gln	Gln	Arg	Leu	His	Arg	Lys	Lys	Gln
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Leu	Glu	Asn	Glu	Met	Met	Arg	Val	Gly	Leu	Ser	Gln	Asp	Ala	Gln	Asp
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Gln	Met	Arg	Lys	Met	Leu	Cys	Gln	Lys	Glu	Ser	Asn	Tyr	Ile	Arg	Leu
		515					520					525			
Lys	Arg	Ala	Lys	Met	Asp	Lys	Ser	Met	Phe	Val	Lys	Ile	Lys	Thr	Leu
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Gly	Ile	Gly	Ala	Phe	Gly	Glu	Val	Cys	Leu	Ala	Arg	Lys	Val	Asp	Thr
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Lys	Ala	Leu	Tyr	Ala	Thr	Lys	Thr	Leu	Arg	Lys	Lys	Asp	Val	Leu	Leu
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Arg	Asn	Gln	Val	Ala	His	Val	Lys	Ala	Glu	Arg	Asp	Ile	Leu	Ala	Glu
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Ala	Asp	Asn	Glu	Trp	Val	Val	Arg	Leu	Tyr	Tyr	Ser	Phe	Gln	Asp	Lys
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Tyr	Ile	Ala	Glu	Leu	Thr	Cys	Ala	Val	Glu	Ser	Val	His	Lys	Met	Gly
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Phe	Ile	His	Arg	Asp	Ile	Lys	Pro	Asp	Asn	Ile	Leu	Ile	Asp	Arg	Asp
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	675						680					685			
Thr	His	Asp	Ser	Lys	Tyr	Tyr	Gln	Ser	Gly	Asp	His	Pro	Arg	Gln	Asp
	690				695					700					
Ser	Met	Asp	Phe	Ser	Asn	Glu	Trp	Gly	Asp	Pro	Ser	Asn	Cys	Arg	Cys
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Arg	Cys	Leu	Ala	His	Ser	Leu	Val	Gly	Thr	Pro	Asn	Tyr	Ile	Ala	Pro
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Glu	Val	Leu	Leu	Arg	Thr	Gly	Tyr	Thr	Gln	Leu	Cys	Asp	Trp	Trp	Ser
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Val	Gly	Val	Ile	Leu	Cys	Glu	Met	Leu	Val	Gly	Gln	Pro	Pro	Phe	Leu
	770					775				780					
Ala	Gln	Thr	Pro	Leu	Glu	Thr	Gln	Met	Lys	Val	Ile	Ile	Trp	Gln	Thr
785					790					795					800
Ser	Leu	His	Ile	Pro	Pro	Gln	Ala	Lys	Leu	Ser	Pro	Glu	Ala	Ser	Asp
				805				810						815	
Leu	Ile	Ile	Lys	Leu	Cys	Arg	Gly	Pro	Glu	Asp	Arg	Leu	Gly	Lys	Asn
			820					825					830		
Gly	Ala	Asp	Glu	Ile	Lys	Ala	His	Pro	Phe	Phe	Lys	Thr	Ile	Asp	Phe
		835					840					845			
Ser	Ser	Asp	Leu	Arg	Gln	Gln	Ser	Ala	Ser	Tyr	Ile	Pro	Lys	Ile	Thr
	850					855				860					
His	Pro	Thr	Asp	Thr	Ser	Asn	Phe	Asp	Pro	Val	Asp	Pro	Asp	Lys	Leu
865					870					875					880
Trp	Ser	Asp	Gly	Ser	Glu	Glu	Glu	Asn	Ile	Ser	Asp	Thr	Leu	Ser	Gly
				885				890						895	
Trp	Tyr	Lys	Asn	Gly	Lys	His	Pro	Glu	His	Ala	Phe	Tyr	Glu	Phe	Thr
			900					905					910		
Phe	Arg	Arg	Phe	Phe	Asp	Asp	Asn	Gly	Tyr	Pro	Tyr	Asn	Tyr	Pro	Lys
		915					920					925			
Pro	Ile	Glu	Tyr	Glu	Tyr	Ile	His	Ser	Gln	Gly	Ser	Glu	Gln	Gln	Ser
	930					935					940				
Asp	Glu	Asp	Asp	Gln	His	Thr	Ser	Ser	Asp	Gly	Asn	Asn	Arg	Asp	Leu
945					950					955					960
Val	Tyr	Val													